

SEQUENCE LISTING

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Ladner, Robert C

<120> BINDING PEPTIDES FOR CARCINOEMBRYONIC ANTIGEN (CEA)

<130> Sequence Listing DYX-016.0 US

<140> (not yet assigned)
<141> 2000-04-03

<160> 107

<170> PatentIn Ver. 2.1

<210> 1
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polypeptide

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<223> Xaa is Asn, Asp or is absent

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<223> Xaa is Trp

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<222> (15)

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<222> (16)

<223> Xaa is Leu, Ser, Trp or Tyr

<400> 1

Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa
1 5 10 15

<210> 2
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preferred CEA binding moieties

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1 5 10 15

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<220>
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Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys
1 5 10

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<220>
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polypeptide

<400> 4
Asn Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asn Ser Tyr
1 5 10 15

<210> 5
<211> 16
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<220>
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<400> 5
Asp Trp Val Cys Glu Asn Lys Lys Asp Gln Trp Thr Cys Asn Leu Leu
1 5 10 15

<210> 6
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<220>
<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 6

Asn Trp Asp Cys Met Phe Gly Ala Glu Gly Trp Ala Cys Ser Pro Trp
1 5 10 15

<210> 7

<211> 16

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<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 7

Asp Trp Val Cys Glu Lys Thr Thr Gly Gly Tyr Val Cys Gln Pro Leu
1 5 10 15

<210> 8

<211> 16

<212> PRT

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<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 8

Asn Trp Phe Cys Glu Met Ile Gly Arg Gln Trp Gly Cys Val Pro Ser
1 5 10 15

<210> 9

<211> 16

<212> PRT

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<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 9

Asp Trp Val Cys Asn Phe Asp Gln Gly Leu Ala His Cys Phe Pro Ser
1 5 10 15

<210> 10

<211> 12
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<220>
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domain for design of microprotein display library

<220>
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<222> (1)..(12)
<223> amino acid positions 4 and 9 are invariant Cys;
all other positions Xaa are varied but not Cys, to
provide a library of 2x10(8) different peptides
based on the template sequence

<400> 10
Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa
1 5 10

<210> 11
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<220>
<223> Description of Artificial Sequence: parental
domain for design of microprotein display library

<220>
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<223> amino acid positions 3 and 9 are invariant Cys;
all other positions Xaa are varied but not Cys, to
provide a library of 1x10(9) different peptides
based on the template sequence

<400> 11
Xaa Xaa Cys Xaa Xaa Xaa Xaa Cys Xaa Xaa
1 5 10

<210> 12
<211> 12
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<220>
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domain for design of microprotein display library

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<223> amino acid positions 3 and 10 are invariant Cys;
all other positions Xaa are varied but not Cys, to
provide a library of 1×10^9 different peptides
based on the template sequence

<400> 12
Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa
1 5 10

<210> 13
<211> 16
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<220>
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domain for design of microprotein display library

<220>
<221> VARIANT
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<223> amino acid positions 4 and 13 are invariant Cys;
all other positions Xaa are varied but not Cys, to
provide a library of 2.5×10^8 different peptides
based on the template sequence

<400> 13
Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa
1 5 10 15

<210> 14
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sublibrary sequence used in designing focused
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<220>
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<400> 14
Xaa Xaa Xaa Cys Xaa Xaa Lys Lys Asp Gln Trp Thr Cys Asn Leu Leu
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secondary library

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1 5 10 15

<210> 16
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secondary library

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<222> (8)..(12)
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1 5 10 15

<210> 18
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<223> Xaa is any amino acid except Cys

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<223> Xaa is any amino acid except Cys

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1 5 10 15

<210> 19
<211> 16
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secondary library

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<223> Xaa is any amino acid except Cys

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<223> Xaa is any amino acid except Cys

<400> 19
Asn Trp Val Cys Xaa Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Ser Tyr
1 5 10 15

<210> 20
<211> 16
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secondary library

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<223> Xaa is any amino acid except Cys

<400> 20
Xaa Trp Xaa Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Xaa Xaa Xaa
1 5 10 15

<210> 21
<211> 16
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<223> Description of Artificial Sequence: isolate of
TN10/9 library found not to bind CEA

<400> 21
Asn Trp Arg Cys Lys Leu Phe Pro Arg Tyr Pro Tyr Cys Ser Ser Trp
1 5 10 15

<210> 22
<211> 15

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: isolate of
TN10/9 library found not to bind CEA

<400> 22

Arg Tyr Cys Glu Phe Phe Pro Trp Ser Leu His Cys Gly Arg Pro
1 5 10 15

<210> 23

<211> 16

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<223> Description of Artificial Sequence: conserved
amino acid positions in first family of CEA
binding peptides

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<223> X is Asn, Leu, Met or Phe

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<221> VARIANT

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<223> X is Asp, Gly, Ile, Lys, Phe or Thr

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<222> (9)

<223> X is Arg, Asn, Asp, Glu or Gly

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<222> (12)

<223> X is Ala, Gly, His, Phe, Thr or Val

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<221> VARIANT

<222> (15)

<223> X is Arg, Leu, Pro or Ser

<400> 23

Asp Trp Val Cys Glu Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Xaa Leu
1 5 10 15

<210> 24
<211> 27
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<220>
<223> Description of Artificial Sequence: synthetic CEA
binding peptide with C-terminal immobilization
sequence

<400> 24
Ser Asn Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asn Ser
1 5 10 15

Tyr Ala Pro Gly Gly Glu Gly Gly Ser Lys
20 25

<210> 25
<211> 27
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<220>
<223> Description of Artificial Sequence: synthetic CEA
binding peptide with C-terminal immobilization
sequence

<400> 25
Ser Asp Trp Val Cys Glu Asn Lys Lys Asp Gln Trp Thr Cys Asn Leu
1 5 10 15

Leu Ala Pro Gly Gly Glu Gly Gly Ser Lys
20 25

<210> 26
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<220>
<223> Description of Artificial Sequence: synthetic CEA
binding peptide with C-terminal immobilization

sequence

<400> 26
Ser Asn Trp Asp Cys Met Phe Gly Ala Glu Gly Trp Ala Cys Ser Pro
1 5 10 15
Trp Ala Pro Gly Gly Glu Gly Gly Ser Lys
20 25

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<211> 27
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: synthetic CEA
binding peptide with C-terminal immobilization
sequence

<400> 27
Ser Asp Trp Val Cys Glu Leu Thr Thr Gly Gly Tyr Val Cys Gln Pro
1 5 10 15
Leu Ala Pro Gly Gly Glu Gly Gly Ser Lys
20 25

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<212> PRT
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<220>
<223> Description of Artificial Sequence: C-terminal
sequence for immobilizing peptides

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Ala Pro Gly Gly Glu Gly Gly Ser Lys
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<210> 29
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<223> Description of Artificial Sequence: template
sequence for sublibrary used in construction of
focused secondary display library

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<223> X is any amino acid except Cys

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Xaa Xaa Xaa Cys Xaa Xaa Lys Lys Asp Gln Trp Thr Cys Asn Leu Leu
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focused secondary display library

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1 5 10 15

<210> 31
<211> 16
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<220>
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focused secondary display library

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1 5 10 15

<210> 33
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1 5 10 15

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focused secondary display library

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<223> X is any amino acid except Cys

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Asn Trp Val Cys Xaa Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Ser Tyr
1 5 10 15

<210> 35

<211> 16

<212> PRT

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<220>

<223> Description of Artificial Sequence: template
sequence for sublibrary used in construction of
focused secondary display library

<220>

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<222> (1)

<223> X is any amino acid except Cys

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<221> VARIANT

<222> (3)

<223> X is any amino acid except Cys

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<222> (14)..(16)

<223> X is any amino acid except Cys

<400> 35

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1 5 10 15

<210> 36

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<212> PRT

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<220>

<223> Description of Artificial Sequence: family of CEA
binding polypeptides

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<221> VARIANT

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<223> Xaa is Asp, Asn, Ala or Ile

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Ser, Val, Trp or Tyr

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or Trp

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<223> Xaa is Gln or Lys

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<223> Xaa is Phe, Thr, Met, Ser, Ala, Asn, Val, His,
Ile, Pro, Trp or Tyr

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Glu, Thr, Lys or Trp

<220>
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<222> (16)
<223> Xaa is Leu, Met, Val, Tyr, Ala, Ile, Trp, His,
Pro, Gln, Glu, Phe, Lys or Arg

<400> 36
Xaa Trp Xaa Cys Xaa Xaa Xaa Xaa Xaa Trp Xaa Cys Xaa Xaa Xaa
1 5 10 15

<210> 37
<211> 16
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 37
Asp Trp Met Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Leu Met
1 5 10 15

<210> 38
<211> 16
<212> PRT
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<220>
<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 38
Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Leu Met
1 5 10 15

<210> 39
<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 39

Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Gln Met
1 5 10 15

<210> 40

<211> 16

<212> PRT

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<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 40

Asn Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Gln Glu
1 5 10 15

<210> 41

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 41

Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Gln Val Lys
1 5 10 15

<210> 42

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 42

Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Val Met
1 5 10 15

<210> 43

<211> 16

<212> PRT

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<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 43

Asp Trp Met Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Gln Ile
1 5 10 15

<210> 44

<211> 16

<212> PRT

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<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 44

Ile Trp Asp Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Pro Ala Pro
1 5 10 15

<210> 45

<211> 16

<212> PRT

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<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 45

Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Ile Arg
1 5 10 15

<210> 46
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<223> Description of Artificial Sequence: CEA binding polypeptide

<400> 46
Asp Trp Met Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Val Val
1 5 10 15

<210> 47
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<400> 47
Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Ala Ile
1 5 10 15

<210> 48
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<223> Description of Artificial Sequence: CEA binding polypeptide

<400> 48
Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Met Ala
1 5 10 15

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<400> 49

Asp Trp Val Cys Glu Phe Leu Lys Met Gln Trp Ala Cys Asn Val Leu
1 5 10 15

<210> 50

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<223> Description of Artificial Sequence: CEA binding polypeptide

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Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asn Val Met
1 5 10 15

<210> 51

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<223> Description of Artificial Sequence: CEA binding polypeptide

<400> 51

Ala Trp Pro Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Pro Pro Gln
1 5 10 15

<210> 52

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